U.S. Patent Application No. 09/706,243

REGULATION OF ENDOGENOUS GENE EXPRESSION IN CELLS USING ZINC FINGER PROTEINS

Pending Claims: October 2001

A method of inhibiting expression of an endogenous cellular gene in a 118. cell, the method comprising the step of:

contacting a first target site in the endogenous cellular gene with an engineered zinc finger protein, wherein the Kd of the zinc finger protein is less than about 25 nM; thereby inhibiting expression of the endogenous cellular gene.

- The method of claim 118, wherein the step of contacting further comprises 119. contacting a second target site in the endogenous cellular gene with a second zinc finger protein.
- The method of claim 119, wherein the first and second target sites are 120. adjacent.
- The method of claim 120, wherein the first and second zinc finger proteins 121. are covalently linked to form a fusion protein.
- The method of claim 118, wherein the first zinc finger protein is a fusion 122. protein comprising a regulatory domain.
- The method of claim 122, wherein the first zinc finger protein is a fusion 123. protein comprising at least two regulatory domains.
- The method of claim 119, wherein the first and second zinc finger proteins 124. are fusion proteins, each comprising a regulatory domain.
- The method of claim 124, wherein the first and second zinc finger protein are fusion proteins, each comprising at least two regulatory domains.

126. A method of inhibiting expression of an endogenous cellular gene in a cell, the method comprising the step of:

contacting a target site in the endogenous cellular gene with an engineered fusion zinc finger protein comprising six fingers and a regulatory domain, wherein the K_d of the zinc finger protein is less than about 25 nM;

thereby inhibiting expression of the endogenous cellular gene.

- 127. The method of claim 118, wherein the cell is selected from the group consisting of an animal cell, a plant cell, a bacterial cell, a protozoal cell, and a fungal cell.
 - 128. The method of claim 127, wherein the cell is a mammalian cell.
 - 129. The method of claim 128, wherein the cell is a human cell.
- 130. The method of claim 118, wherein expression of the endogenous cellular gene is inhibited by at least about 20%.
- 131. The method of claim 118, wherein the endogenous cellular gene is selected from the group consisting of VEGF, ERa, IGF-I, c-myc, c-myb, ICAM, and Her2/Neu.
 - 132. The method of claim 131, wherein the endogenous cellular gene is VEGF.
- 133. The method of claim 118, wherein the inhibition of gene expression prevents gene activation.
- 134. The method of claim 122, wherein the regulatory domain is selected from the group consisting of a transcriptional repressor, an endonuclease, a methyl transferase, and a histone deacetylase.

- 135. The method of claim 124, wherein the regulatory domain is selected from the group consisting of a transcriptional repressor, an endonuclease, a methyl transferase, and a histone deacetylase.
- of first administering to the cell a delivery vehicle comprising the zinc finger protein, wherein the delivery vehicle comprises a liposome or a membrane translocation polypeptide.
- 137. The method of claim 118, wherein the target site is upstream of a transcription initiation site of the endogenous cellular gene.
- 138. The method of claim 118, wherein the target site is adjacent to a transcription initiation site of the endogenous cellular genc.
- 139. The method of claim 118, wherein the zinc finger protein comprises an SP-1 backbone.
- 140. The method of claim 139, wherein the zinc finger protein comprises a regulatory domain and is humanized.
- 141. A method of activating expression of an endogenous cellular gene, the method comprising the step of:

zinc finger protein, wherein the K_d of the zinc finger protein is less than about 25 nM; thereby activating expression of the endogenous cellular gene.

- 142. The method of claim 141, wherein the step of contacting further comprises contacting a second target site in the endogenous cellular gene with a second zinc finger protein.
- 143. The method of claim 142, wherein the first and second target sites are adjacent.

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- 144. The method of claim 143, wherein the first and second zinc finger proteins are covalently linked to form a fusion protein.
- 145. The method of claim 141, wherein the first zinc finger protein is a fusion protein comprising a regulatory domain.
- 146. The method of claim 145, wherein the first zinc finger protein is a fusion protein comprising at least two regulatory domains.
- 147. The method of claim 142, wherein the first and second zinc finger proteins are fusion proteins, each comprising a regulatory domain.
- 148. The method of claim 147, wherein the first and the second zinc finger protein are fusion proteins, each comprising at least two regulatory domains.
- 149. A method of activating expression of an endogenous cellular gene, the method comprising the step of:

contacting a target site in the endogenous cellular gene with an engineered fusion zinc finger protein comprising six fingers and a regulatory domain, wherein the K_d of the zinc finger protein is less than about 25 nM;

thereby activating expression of the endogenous cellular gene.

- 150. The method of claim 141, wherein the cell is selected from the group consisting of an animal cell, a plant cell, a bacterial cell, a protozoal cell, and a fungal cell.
 - 151. The method of claim 150, wherein the cell is a mammalian cell.
 - 152. The method of claim 151, wherein the cell is a human cell.

- 153. The method of claim 141, wherein expression of the endogenous cellular gene is activated to at least about 150%.
- 154. The method of claim 141, wherein the endogenous cellular gene is selected from the group consisting of FAD2-1, EPO, GM-CSF, GDNF, VEGF, and LDL-R.
 - 155. The method of claim 154, wherein the endogenous cellular genc is VEQF.
- 156. The method of claim 141, wherein the activation of gene expression prevents repression of gene expression.
- 157. The method of claim 145, wherein the regulatory domain is selected from the group consisting of a transcriptional activator and a histone acetyltransferase.
- 158. The method of claim 147, wherein the regulatory domain is selected from the group consisting of a transcriptional activator and a histone acetyltransferase.
- 159. The method of claim 141, wherein the target site is upstream of a transcription initiation site of the endogenous cellular gene.
- 160. The method of claim 141, wherein the target site is adjacent to a transcription initiation site of the endogenous cellular gene.
- 161. The method of claim 141, wherein the zinc finger protein comprises an SP-1 backbone.
- 162 The method of claim 161, wherein the zinc finger protein comprises a regulatory domain and is humanized.
- 163. A method of modulating expression of an endogenous cellular gene in a cell, the method comprising the step of:

contacting a first target site in the endogenous cellular gene with an engineered zinc finger protein;

thereby modulating expression of the endogenous cellular gene.

- 164. The method of claim 163, wherein the step of contacting further comprises contacting a second target site in the endogenous cellular gene with a second zinc finger protein.
- 165. The method of claim 164, wherein the first and second target sites are adjacent.
- 166. The method of claim 165, wherein the first and second zinc finger proteins are covalently linked to form a fusion protein.
- 167. The method of claim 163, wherein the first zinc finger protein is a fusion protein comprising a regulatory domain.
- 168. The method of claim 167, wherein the first zinc finger protein is a fusion protein comprising at least two regulatory domains.
- 169. The method of claim 164, wherein the first and second zinc finger proteins are fusion proteins, each comprising a regulatory domain.
- 170. The method of claim 169, wherein the first and second zinc finger protein are fusion proteins, each comprising at least two regulatory domains.
- 171. A method of modulating expression of an endogenous cellular gene in a cell, the method comprising the step of:

contacting a target site in the endogenous cellular gene with an engineered fusion zinc finger protein comprising six fingers and a regulatory domain;

thereby modulating expression of the endogenous cellular gene.

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- 172. The method of claim 163, wherein the cell is selected from the group consisting of an animal cell, a plant cell, a bacterial cell, a protozoal cell, and a fungal cell.
 - 173. The method of claim 172, wherein the cell is a mammalian cell.
 - 174. The method of claim 173, wherein the cell is a human cell.
- 175. The method of claim 163, wherein the endogenous cellular gene is selected from the group consisting of VEGF, ERa, IGF-I, c-myc, c-myb, ICAM, Her2/Neu, FAD2-1, EPO, GM-CSF, GDNF, and LDL-R.
 - 176. The method of claim 175, wherein the endogenous cellular genc is VEGF.
- 177. The method of claim 167, wherein the regulatory domain is selected from the group consisting of a transcriptional repressor, a transcriptional activator, an endonuclease, a methyl transferase, a histone acetyltransferase, and a histone deacetylase.
- 178. The method of claim 169, wherein the regulatory domain is selected from the group consisting of a transcriptional repressor, a transcriptional activator, an endonuclease, a methyl transferase, a histone acetyltransferase, and a histone deacetylase.
- 179. The method of claim 163, wherein the method further comprises the step of first administering to the cell a delivery vehicle comprising the zinc finger protein, wherein the delivery vehicle comprises a liposome or a membrane translocation polypeptide.
- 180. The method of claim 163, wherein the target site is upstream of a transcription initiation site of the endogenous cellular gene.
- 181. The method of claim 163, wherein the target site is adjacent to a transcription initiation site of the endogenous cellular gene.

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182. The method of claim 163, wherein the zinc finger protein comprises an SP-1 backbone.

183. The method of claim 182, wherein the zinc finger protein comprises a regulatory domain and is humanized.